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P#21

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/141,220CDATE: 01/19/2001
TIME: 14:48:27Input Set : A:\Hs1021.app
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3 <110> APPLICANT: Bannon, Gary A
 4 Burks, A Wesley
 5 Sampson, Hugh
 6 Sosin, Howard
 8 <120> TITLE OF INVENTION: Methods and Reagents for Decreasing Clinical Reaction
 to Allergy
 11 <130> FILE REFERENCE: HS1021
 13 <140> CURRENT APPLICATION NUMBER: 09/141,220C
 14 <141> CURRENT FILING DATE: 1998-08-27
 16 <150> PRIOR APPLICATION NUMBER: PCT/US96/15222
 17 <151> PRIOR FILING DATE: 1996-09-23
 19 <150> PRIOR APPLICATION NUMBER: 60/074590
 20 <151> PRIOR FILING DATE: 1998-02-13
 22 <150> PRIOR APPLICATION NUMBER: 60/074624
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 25 <150> PRIOR APPLICATION NUMBER: 60/074633
 26 <151> PRIOR FILING DATE: 1998-02-13
 28 <160> NUMBER OF SEQ ID NOS: 6
 30 <170> SOFTWARE: PatentIn Ver. 2.1
 32 <210> SEQ ID NO: 1
 33 <211> LENGTH: 1930
 34 <212> TYPE: DNA
 35 <213> ORGANISM: Peanut
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 44 ccgtcaaccc cgaaggaggagg aaggaggccg atggggacca gctggacccg gggagcgttg 420
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 60 ccagcttcag gacctggaca tgatgcac ctgtgttagag atcaaagaag gagctttgtt 1380

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62 ctttgaactc gtggctgttaa gaaaagagca acaacaggagg qgacggcgaa aagaagagga 1500
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66 aggtgtataa gacaatgtga tagaccatg agagaagccaa gcaaggatt tagcattcc 1740
67 tgggtgggtt gacaatgtt gaaagctcat caaaaaaaaa aaaaaaccag aaaaaatctc actttgtgaa 1800
68 tgctgtctt caatctcaat ctaatctcc gtgttcttctt gaaaaaaggtt ctccctgagaa 1860
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 200 Lys Thr Glu Asn Pro Cys Ala Gln Arg Cys Leu Gln Ser Cys Gln Gln
 201 35 40 45
 203 Glu Pro Asp Asp Leu Lys Gln Lys Ala Cys Glu Ser Arg Cys Thr Lys
 204 50 55 60
 206 Leu Glu Tyr Asp Pro Arg Leu Val Tyr Asp Pro Arg Gly His Thr Gly
 207 65 70 75 80
 209 Thr Thr Asn Gln Arg Ser Pro Pro Gly Glu Arg Thr Arg Gly Arg Gln
 210 85 90 95
 212 Pro Gly Asp Tyr Asp Asp Asp Arg Arg Gln Pro Arg Arg Glu Glu Gly
 213 100 105 110
 215 Gly Arg Trp Gly Pro Ala Gly Pro Arg Glu Arg Gln Arg Glu Glu Asp
 216 115 120 125
 218 Trp Arg Gln Pro Arg Glu Asp Trp Arg Arg Pro Ser His Gln Gln Pro
 219 130 135 140
 221 Arg Lys Ile Arg Pro Glu Gly Arg Glu Gly Glu Gln Glu Trp Gly Thr
 222 145 150 155 160
 224 Pro Gly Ser His Val Arg Glu Glu Thr Ser Arg Asn Asn Pro Phe Tyr
 225 165 170 175
 227 Phe Pro Ser Arg Arg Phe Ser Thr Arg Tyr Gly Asn Gln Asn Gly Arg
 228 180 185 190
 230 Ile Arg Val Leu Gln Arg Phe Asp Gln Arg Ser Arg Gln Phe Gln Asn
 231 195 200 205
 233 Leu Gln Asn His Arg Ile Val Gln Ile Glu Ala Lys Pro Asn Thr Leu
 234 210 215 220
 236 Val Leu Pro Lys His Ala Asp Ala Asp Asn Ile Leu Val Ile Gln Gln
 237 225 230 235 240
 239 Gly Gln Ala Thr Val Thr Val Ala Asn Gly Asn Asn Arg Lys Ser Phe
 240 245 250 255
 242 Asn Leu Asp Glu Gly His Ala Leu Arg Ile Pro Ser Gly Phe Ile Ser
 243 260 265 270
 245 Tyr Ile Leu Asn Arg His Asp Asn Gln Asn Leu Arg Val Ala Lys Ile
 246 275 280 285
 248 Ser Met Pro Val Asn Thr Pro Gly Gln Phe Glu Asp Phe Phe Pro Ala
 249 290 295 300
 251 Ser Ser Arg Asp Gln Ser Ser Tyr Leu Gln Glu Phe Ser Arg Asn Thr
 252 305 310 315 320

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 257 Leu Glu Glu Asn Ala Gly Gly Glu Gln Glu Glu Arg Gly Gln Arg Arg
 258 340 345 350
 260 Trp Ser Thr Arg Ser Ser Glu Asn Asn Glu Gly Val Ile Val Lys Val
 261 355 360 365
 263 Ser Lys Glu His Val Glu Glu Leu Thr Lys His Ala Lys Ser Val Ser
 264 370 375 380
 266 Lys Lys Gly Ser Glu Glu Gly Asp Ile Thr Asn Pro Ile Asn Leu
 267 385 390 395 400
 269 Arg Glu Gly Glu Pro Asp Leu Ser Asn Asn Phe Gly Lys Leu Phe Glu
 270 405 410 415
 272 Val Lys Pro Asp Lys Lys Asn Pro Gln Leu Gln Asp Leu Asp Met Met
 273 420 425 430
 275 Leu Thr Cys Val Glu Ile Lys Glu Gly Ala Leu Met Leu Pro His Phe
 276 435 440 445
 278 Asn Ser Lys Ala Met Val Ile Val Val Val Asn Lys Gly Thr Gly Asn
 279 450 455 460
 281 Leu Glu Leu Val Ala Val Arg Lys Glu Gln Gln Arg Gly Arg Arg
 282 465 470 475 480
 284 Glu Glu Glu Asp Glu Asp Glu Glu Glu Gly Ser Asn Arg Glu
 285 485 490 495
 287 Val Arg Arg Tyr Thr Ala Arg Leu Lys Glu Gly Asp Val Phe Ile Met
 288 500 505 510
 290 Pro Ala Ala His Pro Val Ala Ile Asn Ala Ser Ser Glu Leu His Leu
 291 515 520 525
 293 Leu Gly Phe Gly Ile Asn Ala Glu Asn Asn His Arg Ile Phe Leu Ala
 294 530 535 540
 296 Gly Asp Lys Asp Asn Val Ile Asp Gln Ile Glu Lys Gln Ala Lys Asp
 297 545 550 555 560
 299 Leu Ala Phe Pro Gly Ser Gly Glu Gln Val Glu Lys Leu Ile Lys Asn
 300 565 570 575
 302 Gln Lys Glu Ser His Phe Val Ser Ala Arg Pro Gln Ser Gln Ser Gln
 303 580 585 590
 305 Ser Pro Ser Ser Pro Glu Lys Glu Ser Pro Glu Lys Glu Asp Gln Glu
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 323 ccctgcgagc aacatctcat gcagaagatc caacgtgacg aggattcata tgaacggac 180
 324 cgtacagcc ctaglcagcc cctagtcctat atgatcgagg aggcgcgtgaa 240

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